

# SEQUENCE LISTING

<110> Yoganathan, Thillainathan  
Delaney, Allen

<120> CAMK-X1 and its Uses

<130> KINE024

<140> Unassigned

<141> 2001-09-20

<160> 13

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 2447

<212> DNA

<213> H. sapiens

<220>

<221> CDS

<222> (70)...(1498)

<400> 1

tggagtggga gctcaagcag gattcttccc gagtccttgg catcctcaga agcttcaact	60
ctggaggca atg ggt cga aag gaa gaa gat gac tgc agt tcc tgg aag aaa	111
Met Gly Arg Lys Glu Glu Asp Asp Cys Ser Ser Trp Lys Lys	
1 5 10	

cag acc acc aac atc cgg aaa acc ttc att ttt atg gaa gtg ctg gga	159
Gln Thr Thr Asn Ile Arg Lys Thr Phe Ile Phe Met Glu Val Leu Gly	
15 20 25 30	

tca gga gct ttc tca gaa gtt ttc ctg gtg aag caa aga ctg act ggg	207
Ser Gly Ala Phe Ser Glu Val Phe Leu Val Lys Gln Arg Leu Thr Gly	
35 40 45	

aag ctc ttt gct ctg aag tgc atc aag aag tca cct gcc ttc cgg gac	255
Lys Leu Phe Ala Leu Lys Cys Ile Lys Lys Ser Pro Ala Phe Arg Asp	
50 55 60	

agc agc ctg gag aat gag att gct gtg ttg aaa aag atc aag cat gaa	303
Ser Ser Leu Glu Asn Glu Ile Ala Val Leu Lys Lys Ile Lys His Glu	
65 70 75	

aac att gtg acc ctg gag gac atc tat gag agc acc acc cac tac tac	351
Asn Ile Val Thr Leu Glu Asp Ile Tyr Glu Ser Thr Thr His Tyr Tyr	
80 85 90	

ctg gtc atg cag ctt gtt tct ggt ggg gag ctc ttt gac cgg atc ctg	399
---	-----

Leu	Val	Met	Gln	Leu	Val	Ser	Gly	Gly	Glu	Leu	Phe	Asp	Arg	Ile	Leu	
95					100					105					110	
gag	cgg	ggg	gtc	tac	aca	gag	aag	gat	gcc	agt	ctg	gtg	atc	cag	cag	447
Glu	Arg	Gly	Val	Tyr	Thr	Glu	Lys	Asp	Ala	Ser	Leu	Val	Ile	Gln	Gln	
				115					120					125		
gtc	ttg	tcg	gca	gtg	aaa	tac	cta	cat	gag	aat	ggc	atc	gtc	cac	aga	495
Val	Leu	Ser	Ala	Val	Lys	Tyr	Leu	His	Glu	Asn	Gly	Ile	Val	His	Arg	
			130					135					140			
gac	tta	aag	ccc	gaa	aac	ctg	ctt	tac	ctt	acc	cct	gaa	gag	aac	tct	543
Asp	Leu	Lys	Pro	Glu	Asn	Leu	Leu	Tyr	Leu	Thr	Pro	Glu	Glu	Asn	Ser	
		145					150					155				
aag	atc	atg	atc	act	gac	ttt	ggg	ctg	tcc	aag	atg	gaa	cag	aat	ggc	591
Lys	Ile	Met	Ile	Thr	Asp	Phe	Gly	Leu	Ser	Lys	Met	Glu	Gln	Asn	Gly	
	160					165				170						
atc	atg	tcc	act	gcc	tgt	ggg	acc	cca	ggc	tac	gtg	gct	cca	gaa	gtg	639
Ile	Met	Ser	Thr	Ala	Cys	Gly	Thr	Pro	Gly	Tyr	Val	Ala	Pro	Glu	Val	
	175				180				185					190		
ctg	gcc	cag	aaa	ccc	tac	agc	aag	gct	gtg	gat	tgc	tgg	tcc	atc	ggc	687
Leu	Ala	Gln	Lys	Pro	Tyr	Ser	Lys	Ala	Val	Asp	Cys	Trp	Ser	Ile	Gly	
				195				200						205		
gtc	atc	acc	tac	ata	ttg	ctc	tgt	gga	tac	ccc	ccg	ttc	tat	gaa	gaa	735
Val	Ile	Thr	Tyr	Ile	Leu	Leu	Cys	Gly	Tyr	Pro	Pro	Phe	Tyr	Glu	Glu	
			210					215					220			
acg	gag	tct	aag	ctt	ttc	gag	aag	atc	aag	gag	ggc	tac	tat	gag	ttt	783
Thr	Glu	Ser	Lys	Leu	Phe	Glu	Lys	Ile	Lys	Glu	Gly	Tyr	Tyr	Glu	Phe	
		225					230					235				
gag	tct	cca	ttc	tgg	gat	gac	att	tct	gag	tca	gcc	aag	gac	ttt	att	831
Glu	Ser	Pro	Phe	Trp	Asp	Asp	Ile	Ser	Glu	Ser	Ala	Lys	Asp	Phe	Ile	
		240				245					250					
tgc	cac	ttg	ctt	gag	aag	gat	ccg	aac	gag	cgg	tac	acc	tgt	gag	aag	879
Cys	His	Leu	Leu	Glu	Lys	Asp	Pro	Asn	Glu	Arg	Tyr	Thr	Cys	Glu	Lys	
	255				260				265					270		
gcc	ttg	agt	cat	ccc	tgg	att	gac	gga	aac	acg	gcc	ctc	cac	cgg	gac	927
Ala	Leu	Ser	His	Pro	Trp	Ile	Asp	Gly	Asn	Thr	Ala	Leu	His	Arg	Asp	
				275				280						285		
atc	tac	cca	tca	gtc	agc	ctc	cag	atc	cag	aag	aac	ttt	gct	aag	agc	975
Ile	Tyr	Pro	Ser	Val	Ser	Leu	Gln	Ile	Gln	Lys	Asn	Phe	Ala	Lys	Ser	
			290					295					300			
aag	tgg	agg	caa	gcc	ttc	aac	gca	gca	gct	gtg	gtg	cac	cac	atg	agg	1023
Lys	Trp	Arg	Gln	Ala	Phe	Asn	Ala	Ala	Ala	Val	Val	His	His	Met	Arg	
		305					310					315				

aag	cta	cac	atg	aac	ctg	cac	agc	ccg	ggc	gtc	cgc	cca	gag	gtg	gag		1071
Lys	Leu	His	Met	Asn	Leu	His	Ser	Pro	Gly	Val	Arg	Pro	Glu	Val	Glu		
	320					325					330						
aac	agg	ccg	cct	gaa	act	caa	gcc	tca	gaa	acc	tct	aga	ccc	agc	tcc		1119
Asn	Arg	Pro	Pro	Glu	Thr	Gln	Ala	Ser	Glu	Thr	Ser	Arg	Pro	Ser	Ser		
335					340					345					350		
cct	gag	atc	acc	atc	acc	gag	gca	cct	gtc	ctg	gac	cac	agt	gta	gca		1167
Pro	Glu	Ile	Thr	Ile	Thr	Glu	Ala	Pro	Val	Leu	Asp	His	Ser	Val	Ala		
				355					360					365			
ctc	cct	gcc	ctg	acc	caa	tta	ccc	tgc	cag	cat	ggc	cgc	cgg	ccc	act		1215
Leu	Pro	Ala	Leu	Thr	Gln	Leu	Pro	Cys	Gln	His	Gly	Arg	Arg	Pro	Thr		
			370					375					380				
gcc	cct	ggg	ggc	agg	tcc	ctc	aac	tgc	ctg	gtc	aat	ggc	tcc	ctc	cac		1263
Ala	Pro	Gly	Gly	Arg	Ser	Leu	Asn	Cys	Leu	Val	Asn	Gly	Ser	Leu	His		
		385					390					395					
atc	agc	agc	agc	ctg	gtg	ccc	atg	cat	cag	ggg	tcc	ctg	gcc	gcc	ggg		1311
Ile	Ser	Ser	Ser	Leu	Val	Pro	Met	His	Gln	Gly	Ser	Leu	Ala	Ala	Gly		
	400					405					410						
ccc	tgt	ggc	tgc	tgc	tcc	agc	tgc	ctg	aac	att	ggg	agc	aaa	gga	aag		1359
Pro	Cys	Gly	Cys	Cys	Ser	Ser	Cys	Leu	Asn	Ile	Gly	Ser	Lys	Gly	Lys		
415					420					425					430		
tcc	tcc	tac	tgc	tct	gag	ccc	aca	ctc	ctc	aaa	aag	gcc	aac	aaa	aaa		1407
Ser	Ser	Tyr	Cys	Ser	Glu	Pro	Thr	Leu	Leu	Lys	Lys	Ala	Asn	Lys	Lys		
				435					440					445			
cag	aac	ttc	aag	tcg	gag	gtc	atg	gta	cca	gtt	aaa	gcc	agt	ggc	agc		1455
Gln	Asn	Phe	Lys	Ser	Glu	Val	Met	Val	Pro	Val	Lys	Ala	Ser	Gly	Ser		
			450					455					460				
tcc	cac	tgc	cgg	gca	ggg	cag	act	gga	gtc	tgt	ctc	att	atg	t			1498
Ser	His	Cys	Arg	Ala	Gly	Gln	Thr	Gly	Val	Cys	Leu	Ile	Met				
		465					470					475					
gattcctgga	gcctgtgcct			atgtcactgc			aattttcagg			agacatatte			aactcctctg			1558	
ctcttccaaa	cctgggtgtct			atccggcaga			gggaggaagg			cagagcaagt			ggagcagggc			1618	
ttagcaggag	cagtttctgg			ccagaagcac			cagcctgctg			ccagcggggc			agccccctcat			1678	
aggaggccca	ggagggagcc			ccaaggcgta			gaagccttgt			tgaagctgtg							

attttaatgt ctgccaggag ttctaatect gcctctgttc ccttttctct ccttgaaagt 2338  
 ccagcacacc attcttgatc ttccccagtt tctctgcctt ccaccctcc agcttcacgc 2398  
 tcagtgtgtg gcttaataaaa atggacatat ttttctctaa aaaaaaaaaa 2447

<210> 2  
 <211> 476  
 <212> PRT  
 <213> H. sapiens

<400> 2  
 Met Gly Arg Lys Glu Glu Asp Asp Cys Ser Ser Trp Lys Lys Gln Thr  
 1 5 10 15  
 Thr Asn Ile Arg Lys Thr Phe Ile Phe Met Glu Val Leu Gly Ser Gly  
 20 25 30  
 Ala Phe Ser Glu Val Phe Leu Val Lys Gln Arg Leu Thr Gly Lys Leu  
 35 40 45  
 Phe Ala Leu Lys Cys Ile Lys Lys Ser Pro Ala Phe Arg Asp Ser Ser  
 50 55 60  
 Leu Glu Asn Glu Ile Ala Val Leu Lys Lys Ile Lys His Glu Asn Ile  
 65 70 75 80  
 Val Thr Leu Glu Asp Ile Tyr Glu Ser Thr Thr His Tyr Tyr Leu Val  
 85 90 95  
 Met Gln Leu Val Ser Gly Gly Glu Leu Phe Asp Arg Ile Leu Glu Arg  
 100 105 110  
 Gly Val Tyr Thr Glu Lys Asp Ala Ser Leu Val Ile Gln Gln Val Leu  
 115 120 125  
 Ser Ala Val Lys Tyr Leu His Glu Asn Gly Ile Val His Arg Asp Leu  
 130 135 140  
 Lys Pro Glu Asn Leu Leu Tyr Leu Thr Pro Glu Glu Asn Ser Lys Ile  
 145 150 155 160  
 Met Ile Thr Asp Phe Gly Leu Ser Lys Met Glu Gln Asn Gly Ile Met  
 165 170 175  
 Ser Thr Ala Cys Gly Thr Pro Gly Tyr Val Ala Pro Glu Val Leu Ala  
 180 185 190  
 Gln Lys Pro Tyr Ser Lys Ala Val Asp Cys Trp Ser Ile Gly Val Ile  
 195 200 205  
 Thr Tyr Ile Leu Leu Cys Gly Tyr Pro Pro Phe Tyr Glu Glu Thr Glu  
 210 215 220  
 Ser Lys Leu Phe Glu Lys Ile Lys Glu Gly Tyr Tyr Glu Phe Glu Ser  
 225 230 235 240  
 Pro Phe Trp Asp Asp Ile Ser Glu Ser Ala Lys Asp Phe Ile Cys His  
 245 250 255  
 Leu Leu Glu Lys Asp Pro Asn Glu Arg Tyr Thr Cys Glu Lys Ala Leu  
 260 265 270  
 Ser His Pro Trp Ile Asp Gly Asn Thr Ala Leu His Arg Asp Ile Tyr  
 275 280 285  
 Pro Ser Val Ser Leu Gln Ile Gln Lys Asn Phe Ala Lys Ser Lys Trp  
 290 295 300  
 Arg Gln Ala Phe Asn Ala Ala Val Val His His Met Arg Lys Leu  
 305 310 315 320  
 His Met Asn Leu His Ser Pro Gly Val Arg Pro Glu Val Glu Asn Arg  
 325 330 335  
 Pro Pro Glu Thr Gln Ala Ser Glu Thr Ser Arg Pro Ser Ser Pro Glu  
 340 345 350

Ile	Thr	Ile	Thr	Glu	Ala	Pro	Val	Leu	Asp	His	Ser	Val	Ala	Leu	Pro
		355					360					365			
Ala	Leu	Thr	Gln	Leu	Pro	Cys	Gln	His	Gly	Arg	Arg	Pro	Thr	Ala	Pro
		370				375					380				
Gly	Gly	Arg	Ser	Leu	Asn	Cys	Leu	Val	Asn	Gly	Ser	Leu	His	Ile	Ser
385					390					395					400
Ser	Ser	Leu	Val	Pro	Met	His	Gln	Gly	Ser	Leu	Ala	Ala	Gly	Pro	Cys
				405					410					415	
Gly	Cys	Cys	Ser	Ser	Cys	Leu	Asn	Ile	Gly	Ser	Lys	Gly	Lys	Ser	Ser
			420					425					430		
Tyr	Cys	Ser	Glu	Pro	Thr	Leu	Leu	Lys	Lys	Ala	Asn	Lys	Lys	Gln	Asn
		435					440					445			
Phe	Lys	Ser	Glu	Val	Met	Val	Pro	Val	Lys	Ala	Ser	Gly	Ser	Ser	His
	450					455					460				
Cys	Arg	Ala	Gly	Gln	Thr	Gly	Val	Cys	Leu	Ile	Met				
465					470					475					

<210> 3  
 <211> 25  
 <212> DNA  
 <213> H. sapiens

<400> 3  
 gtggagggcg aggaaactgg ggaag

25

<210> 4  
 <211> 23  
 <212> DNA  
 <213> Homo sapiens

<400> 4  
 ggagggcgag gaaactgggg aag

23

<210> 5  
 <211> 25  
 <212> DNA  
 <213> Homo sapiens

<400> 5  
 gtggagggcg aggaaactgg ggaag

25

<210> 6  
 <211> 31  
 <212> DNA  
 <213> Homo sapiens

<400> 6  
 ctcgagtcac ataatgagac agactccagt c

31

<210> 7  
 <211> 13  
 <212> PRT  
 <213> Homo sapiens

Category	Sub-category	Value
1. General Information	1.1. Name	John Doe
	1.2. Age	35
	1.3. Gender	Male
	1.4. Date of Birth	1988-05-15
	1.5. Address	123 Main St, New York, NY 10001
	1.6. Phone Number	(212) 555-1234
	1.7. Email Address	john.doe@example.com
	1.8. Occupation	Software Engineer
	1.9. Education	B.S. in Computer Science
	1.10. Marital Status	Single
2. Employment History	2.1. Company Name	ABC Corporation
	2.2. Job Title	Senior Software Engineer
	2.3. Start Date	2015-01-01
	2.4. End Date	2020-12-31
	2.5. Salary	\$120,000
	2.6. Reason for Leaving	Seeking new challenges
	2.7. Supervisor	John Smith
	2.8. Performance Rating	Excellent
	2.9. Skills Acquired	Python, JavaScript, React
	2.10. References	John Smith, Manager
3. Education Details	3.1. Institution Name	XYZ University
	3.2. Degree Program	B.S. in Computer Science
	3.3. Graduation Year	2012
	3.4. GPA	3.8
	3.5. Thesis Topic	Advanced Algorithms
	3.6. Advisor	Prof. Jane Doe
	3.7. Honors/Awards	Dean's List
	3.8. Extracurricular Activities	Chess, Coding
	3.9. Contact Information	123 Main St, New York, NY 10001
	3.10. References	Prof. Jane Doe
4. Skills and Interests	4.1. Programming Languages	Python, JavaScript, Java
	4.2. Frameworks/Libraries	React, Django, Spring
	4.3. Tools	Git, Docker, Kubernetes
	4.4. Soft Skills	Teamwork, Communication
	4.5. Languages	English, Spanish
	4.6. Hobbies	Reading, Traveling
	4.7. Certifications	AWS Certified Solutions Architect
	4.8. Projects	Web Application Development
	4.9. Publications	None
	4.10. References	None
5. References	5.1. Name	John Smith
	5.2. Title	Manager
	5.3. Company	ABC Corporation
	5.4. Contact Info	(212) 555-5678
	5.5. Email	john.smith@abc.com
	5.6. Address	456 Main St, New York, NY 10001
	5.7. Date	2021-01-15
	5.8. Signature	[Signature]
	5.9. Stamp	[Stamp]
	5.10. Notes	John is a highly motivated and skilled professional.

```
<210> 8
<211> 15
<212> PRT
<213> Homo sapiens
```

Pro Leu Ala Arg Thr Leu Ser Val Ala Gly Leu Pro Gly Lys Lys  
1 5 10 15

```
<210> 9
<211> 10
<212> PRT
<213> Homo sapiens
```

Pro Leu Ser Arg Thr Leu Ser Val Ser Ser  
1 5 10

```
<210> 10
<211> 30
<212> DNA
<213> Homo sapiens
```

```
<400> 10
gaattcaatg ggtcgaaagg aagaagatga 30
```

```
<210> 11
<211> 31
<212> DNA
<213> Homo sapiens
```

```
<400> 11
ctcgagtcac ataatgagac agactccagt c 31
```

```
<210> 12
<211> 30
<212> DNA
<213> Homo sapiens
```

<400> 12  
gaattcaatg ggtcgaaagg aagaagatga 30

```
<210> 13
<211> 30
<212> DNA
<213> Homo sapiens
```

<400> 13

ctcgagctgg atctggagggc tgactgatgg

30

ctcgagctgg atctggagggc tgactgatgg